Amendments to the Claims

Please amend claims 1 and 25-28 as follows. All pending claims, whether or not amended, are presented below for the Examiner's convenience and will replace all prior versions, and listings, of claims in the application.

1. (Currently amended) A nucleic acid encoding a modified polypeptide with an improved *in vivo* half-life, said modified polypeptide comprising an Ig constant domain or Ig-like constant domain and a salvage receptor binding epitope within said Ig constant domain or Ig-like constant domain, wherein said epitope is absent from the unmodified polypeptide, wherein said salvage receptor binding epitope is taken from ~~a~~ at least one loop of the CH_2 domain of an Fc region of an Ig molecule and wherein said polypeptide in modified form does not comprise an intact CH_2 domain or an intact Fc region.

21. (Previously added) The nucleic acid of claim 1 wherein the Ig domain or Ig-like domain comprises a CH_1 domain.

22. (Previously added) The nucleic acid of claim 1 wherein the unmodified polypeptide is an Fab, an $(Fab)_2$, or a receptor.

23. (Previously added) The nucleic acid of claim 22 wherein the unmodified polypeptide is an anti-CD18 Fab or an anti-CD18 $(Fab)_2$.

24. (Previously added) The nucleic acid of claim 23 wherein the modified polypeptide is human or humanized.

25. (Currently amended) The nucleic acid of claim 1 wherein said salvage receptor epitope binding epitope comprises amino acids from 1 through ~~about~~ 11 of SEQ ID NO: 3.

26. (Currently amended) The nucleic acid of claim 1 wherein said salvage receptor binding epitope comprises amino acids from 1 through ~~about~~ 11 of SEQ ID NO: 3 and amino acids from 1 through ~~about~~ 7 of SEQ ID NO: 11.

27. (Currently amended) The nucleic acid of claim 1 wherein said salvage receptor binding epitope comprises amino acids from 1 through ~~about~~ 11 of SEQ ID NO: 3 and amino acids from 1 through ~~about~~ 8 of SEQ ID NO: 1.

28. (Currently amended) The nucleic acid of claim 1 wherein said salvage receptor comprises amino acids from 1 through ~~about~~ 11 of SEQ ID NO: 3 and amino acids from 1 through ~~about~~ 8 of SEQ ID NO: 31.